	CRITTORS Corrected by the STIC Systems Branch	2
N	lumber: 08/862,6/0 CRF Processing Date: 1/39/97	_
	Changed a file from non-ASCII to ASCII  Verified by:   Verified by:	) :
	Changed the margins in cases where the sequence text was "wrapped down to the next line.	· }
	Edited a format error in the Current Application Data section, specifically:	•
	Edited the Current Application Data section with the actual current number inputted by the applicant was the prior application data; or other	Э
	Added the mandatory heading and subheadings for "Current Application Data".	
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an intege	
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:	
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:	
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:	
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.	
	Inserted colons after headings/subheadings. Headings edited included:	
	Deleted extra, invalid, headings used by an applicant, specifically:	
	Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end o ☐ page numbers throughout text; ☐ other invalid text, such as	– I fi –
	Inserted mandatory headings, specifically:	
	Corrected an obvious error in the response, specifically:	
•	Edited identifiers where upper case is used but lower case is required, or vice versa.	
	Corrected an error in the Number of Sequences field, specifically:	
_	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.	
	Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (endue to a PatentIn bug). Sequences corrected:	·01
	Other: Jeg 6 - corneled (A) LENGTH: respense  - insided Land releven after 1st puellie and Rold	
	Manual Control of the	_

Fram 8

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

#### RAW SEOUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 13:59:47

```
This Raw Listing contains the General Information Section and up to the first 5 pages.

SEQUENCE LISTING

Per, David H. Lloyd

The Sequencing OCT
 1
 2
             General Information:
 3
     (1)
 4
     (i) APPLICANT: Glenn Albrecht, Sydney Brenner, David H. Lloyd
 5
 6
 7
     (ii) TITLE OF INVENTION: Massively Parallel Signature Sequencing
 8
        by Ligation of Encoded Adaptors
 9
     (iii) NUMBER OF SEQUENCES:
10
11
12
     (iv) CORRESPONDENCE ADDRESS:
13
     (A) ADDRESSEE: Stephen C. Macevicz, Lynx Therapeutics, Inc.
     (B) STREET: 3832 Bay Center Place
14
15
     (C) CITY: Hayward
     (D) STATE: California
16
     (E) COUNTRY: USA
17
18
     (F) ZIP: 94545
19
     (V) COMPUTER READABLE FORM:
20
21
     (A) MEDIUM TYPE: 3.5 inch diskette
22
     (B) COMPUTER: IBM compatible
     (C) OPERATING SYSTEM: Windows 3.1/DOS 5.0
23
24
     (D) SOFTWARE: Microsoft Word for Windows, vers. 2.0
25
26
     (vi) CURRENT APPLICATION DATA:
27
     (A) APPLICATION NUMBER:
28
     (B) FILING DATE:
29
     (C) CLASSIFICATION:
30
31
     (vii) PRIOR APPLICATION DATA:
     (A) APPLICATION NUMBER: 08/689,587
32
33
     (B) FILING DATE: 12-AUG-96
34
35
     (vii) PRIOR APPLICATION DATA:
     (A) APPLICATION NUMBER: 08/659,453
36
     (B) FILING DATE: 06-JUN-96
37
38
39
     (viii) ATTORNEY/AGENT INFORMATION:
40
     (A) NAME: Stephen C. Macevicz
     (B) REGISTRATION NUMBER: 30,285
41
42
     (C) REFERENCE/DQCKET NUMBER: 808-lus
43
     (ix) TELECOMMUNICATION INFORMATION:
44
     (A) TELEPHONE: (510) 670-9365
45
     (B) TELEFAX: (510)670-9302
```

### RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 13:59:51

INPUT SET: S19282.raw

```
48
49
     (2) INFORMATION FOR SEQ ID NO: 1:
50
51
     (i) SEQUENCE CHARACTERISTICS:
52
     (A) LENGTH: 28 nucleotides
     (B) TYPE: nucleic acid
53
54
     (C) STRANDEDNESS: double
55
     (D) TOPOLOGY: linear
56
57
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
58
59
    TGGATTCTAG AGAGAGAGA AGAGAGAG
                                                                   28
60
61
     (2) INFORMATION FOR SEQ ID NO: 2:
62
63
     (i) SEQUENCE CHARACTERISTICS:
64
     (A) LENGTH: 16 nucleotides
65
     (B) TYPE: nucleic acid
66
67
     (C) STRANDEDNESS: double
     (D) TOPOLOGY: linear
68
69
70
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
71
72
73
    NNGGATGNNN NNNNNN
                                                                   16
74
75
    (2) INFORMATION FOR SEQ ID NO: 3:
76
77
78
    (i) SEQUENCE CHARACTERISTICS:
79
    (A) LENGTH: 11 nucleotides
    (B) TYPE: nucleic acid
81
     (C) STRANDEDNESS: single
    (D) TOPOLOGY: linear
82
83
84
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
85
86
87
    NRRGATCYNN N
                                                                   11
88
89
90
    (2) INFORMATION FOR SEQ ID NO: 4:
91
92
     (i) SEQUENCE CHARACTERISTICS:
93
     (A) LENGTH: 20 nucleotides
94
    (B) TYPE: nucleic acid
95
     (C) STRANDEDNESS: double
96
    (D) TOPOLOGY: linear
97
98
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
99
```

### RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 13:59:54

INPUT SET: S19282, raw AGTGGCTGGG CATCGGACCG (2) INFORMATION FOR SEQ ID NO: 5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 nucleotides (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5: GGGGCCCAGT CAGCGTCGAT (2) INFORMATION FOR SEQ ID NO: 6: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 70 nucleotides (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6: AAAAGGAGGA GGCCTTGATA GAGAGGACCT GTTTAAACGT TTAAACGGAT CCTCTTCCTC TTCCTCTTCC (2) INFORMATION FOR SEQ ID NO: 7: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 nucleotides (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7: CTAAACCATT GGTATGGGCC AGTGAATTGT AATA (2) INFORMATION FOR SEQ ID NO: 8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 55 nucleotides (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8: CGCGCAGCCC GCATCGTTTA TGCTACAGAC TGTCAGTGCA 

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 13:59:58

INPUT SET: S19282.raw

153	GCTCTCCGAT CCAAA	55
154		
155	(2) INFORMATION FOR SEQ ID NO: 9:	
156	· ·	
157	(i) SEQUENCE CHARACTERISTICS:	
158	(A) LENGTH: 16 nucleotides	
159	(B) TYPE: nucleic acid	
160	(C) STRANDEDNESS: double	
161	(D) TOPOLOGY: linear	
162		
163	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:	
164		
165	TCGATGGGCT AGATTT	16
166		
167		
168	(2) INFORMATION FOR SEQ ID NO: 10:	
169		
170	(i) SEQUENCE CHARACTERISTICS:	
171	(A) LENGTH: 30 nucleotides	
172	(B) TYPE: nucleic acid	
173	(C) STRANDEDNESS: double	
174	(D) TOPOLOGY: linear	
175		
176	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:	
177		
178	ANNNTACAGC TGCATCCCTT GGCGCTGAGG	30
179	/A\ TW70D\/BTAY FOR GRA TO WA 11	
180	(2) INFORMATION FOR SEQ ID NO: 11:	
181	/il GROUENGE GUADAGMEDIGMIGG.	
182	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 30 nucleotides	
184 185	(B) TYPE: nucleic acid (C) STRANDEDNESS: double	
186	(D) TOPOLOGY: linear	
187	(D) TOPOLOGI: IIMeal	
188	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:	
189	(XI) SEQUENCE DESCRIPTION. SEQ ID NO. 11.	
190	NANNTACAGC TGCATCCCTG GGCCTGTAAG	30
191	MANUTACAGE IGENIECETO GGEETGIANG	30
192		
193	(2) INFORMATION FOR SEQ ID NO: 12:	
194	(2) INICIMATION TON DEG ID NO. 12.	
195	(i) SEQUENCE CHARACTERISTICS:	
196	(A) LENGTH: 30 nucleotides	
197	(B) TYPE: nucleic acid	
198	(C) STRANDEDNESS: double	
199	(D) TOPOLOGY: linear	
200		
201	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:	
202	//	
203	CNNNTACAGC TGCATCCCTT GACGGGTCTC	30
204	•	
205	(2) INFORMATION FOR SEQ ID NO: 13:	
	· · · · · · · · · · · · · · · · · · ·	

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 14:00:01

INPUT SET: S19282.raw

		INPU	T SET: S19282.raw
206			
207	(i) SEQUENCE CHARACTERISTICS:		
208	(A) LENGTH: 30 nucleotides		
209	(B) TYPE: nucleic acid		
210	(C) STRANDEDNESS: double		
211	(D) TOPOLOGY: linear		
212	/		
213	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:	13:	
214	NONNERGROOM COCCORDA COCCORDA COCCO		20
215 216	NCNNTACAGC TGCATCCCTG CCCGCACAGT		30
217	(2) INFORMATION FOR SHOUTD NO. 14.		
217	(2) INFORMATION FOR SEQ ID NO: 14:		
219	(i) SEQUENCE CHARACTERISTICS:		
220	(A) LENGTH: 30 nucleotides		
221	(B) TYPE: nucleic acid		
222	(C) STRANDEDNESS: double		
223	(D) TOPOLOGY: linear		
224	(b) Torozoor. Trincar		
225	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:	14:	
226	(, <u>-</u>		
227	GNNNTACAGC TGCATCCCTT CGCCTCGGAC		30
228			
229	(2) INFORMATION FOR SEQ ID NO: 15:		
230			
231	(i) SEQUENCE CHARACTERISTICS:		
232	(A) LENGTH: 30 nucleotides		
233	(B) TYPE: nucleic acid		
234	(C) STRANDEDNESS: double		
235	(D) TOPOLOGY: linear		
236			
237	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:	15:	
238	V6W-01-61-66 - 0561-056-05-1-6-		
239	NGNNTACAGC TGCATCCCTG ATCCGCTAGC		30
240 241			
241	(2) INFORMATION FOR CEO ID NO. 16.		
242	(2) INFORMATION FOR SEQ ID NO: 16:		
244	(i) SEOUENCE CHARACTERISTICS:		
245	(A) LENGTH: 30 nucleotides		·
	(B) TYPE: nucleic acid		
247	(C) STRANDEDNESS: double		
248	(D) TOPOLOGY: linear		
249	(2) 101020011 21.1001		
250	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:	16:	
251	, ,		
<b>₽</b> 252	TNNNTACAGC TGCATCCCTT CCGAACCCGC		30
253			
254	(2) INFORMATION FOR SEQ ID NO: 17:		
255	- · ·		
256	(i) SEQUENCE CHARACTERISTICS:		
257	(A) LENGTH: 30 nucleotides		
258	(B) TYPE: nucleic acid		

### RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 14:00:05

INPUT SET: S19282.raw

#### \*\*\*\*\* PREVIOUSLY ERRORED SEQUENCES - EDITED \*\*\*\*\*

.15	(2) INFORMATION FOR SEQ ID NO: 6:	
.16		
.17	(i) SEQUENCE CHARACTERISTICS:	
.18	(A) LENGTH: 70 nucleotides	
.19	(B) TYPE: nucleic acid	
.20	(C) STRANDEDNESS: double	
21	(D) TOPOLOGY: linear	
.22		
.23	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
24		
.25	AAAAGGAGGA GGCCTTGATA GAGAGGACCT GTTTAAACGT TTAAACGGAT	50
.26	CCTCTTCCTC TTCCTCTTCC	70
.27		
.28		

### **SEQUENCE VERIFICATION REPORT** PATENT APPLICATION *US/08/862,610*

DATE: 07/30/97 TIME: 14:00:06

INPUT SET: S19282.raw

Line

Error

Original Text





#### RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 10:31:16

INPUT SET: S19282.raw

This Raw Listing contains the General **Information Section and those Sequences** containing ERRORS.

```
ices

Corrected Does Not Comply
Needed Needed
                                       SEQUENCE LISTING
1
 2
 3
     (1)
            General Information:
     (i) APPLICANT: Glenn Albrecht, Sydney Brenner, David H. Lloyd
 5
 6
     (ii) TITLE OF INVENTION: Massively Parallel Signature Sequencing by Ligation of En
 7
 8
 9
     (iii) NUMBER OF SEQUENCES:
10
11
     (iv) CORRESPONDENCE ADDRESS:
     (A) ADDRESSEE: Stephen C. Macevicz, Lynx Therapeutics, Inc.
12
     (B) STREET: 3832 Bay Center Place
13
14
     (C) CITY: Hayward
15
     (D) STATE: California
16
     (E) COUNTRY: USA
17
     (F) ZIP: 94545
18
19
     (V) COMPUTER READABLE FORM:
     (A) MEDIUM TYPE: 3.5 inch diskette
20
21
     (B) COMPUTER: IBM compatible
     (C) OPERATING SYSTEM: Windows 3.1/DOS 5.0
22
     (D) SOFTWARE: Microsoft Word for Windows, vers. 2.0
23
24
25
     (vi) CURRENT APPLICATION DATA:
     (A) APPLICATION NUMBER:
26
27
     (B) FILING DATE:
28
     (C) CLASSIFICATION:
29
     (vii) PRIOR APPLICATION DATA:
30
     (A) APPLICATION NUMBER: 08/689,587
31
32
     (B) FILING DATE: 12-AUG-96
33
     (vii) PRIOR APPLICATION DATA:
35
     (A) APPLICATION NUMBER: 08/659,453
36
     (B) FILING DATE: 06-JUN-96
37
38
     (viii) ATTORNEY/AGENT INFORMATION:
39
     (A) NAME > Stephen C. Macevicz
40
     (B) REGISTRATION NUMBER: 30,285
     (C) AREFERENCE/DOCKET NUMBER: 808-lus
41
42
43
     (ix TELECOMMUNICATION INFORMATION:
     (A) TELEPHONE: (510) 670-9365
44
```

(B) TELEFAX: (510)670-9302

45

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 10:31:18

INPUT SET: S19282.raw

46 47

### ERRORED SEQUENCES FOLLOW:

	114	(2) INFORMATION FOR SEQ ID NO: 6:	
	115		
	116	(i) SEQUENCE CHARACTERISTICS:	
>	117	(A) LENGTH: (20) nucleotides To show	
	118	(B) TYPE: nucleic acid	•
	119	(C) STRANDEDNESS: double	
	120	(D) TOPOLOGY: linear	
	121		
	122	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	insert hard return
	123		LIMS OF THE PROPERTY.
	124	AAAAGGAGGA GGCCTTGATA GAGAGGACCT GTTTAAACGT TTAAACGGAT	50 CCTCTTCCTC TTCCTCTTC
	125		
	126		



### SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/862,610

DATE: 07/30/97 TIME: 10:31:19

INPUT SET: S19282.raw

Line

Error

Original Text

117

Entered (20) and Calc. Seq. Length (0) differ

(A) LENGTH: 20 nucleotides